



Figure 2-1

AccI		GT'mk_AC (0 Err) - 2 Fragment(s)
	243	909
Apal		G_GGCC'C (0 Err) - 2 Fragment(s)
	402	750
BamHI		G'GATC_C (0 Err) - 2 Fragment(s)
	200	952
EcoRI		G'AATT_C (0 Err) - 2 Fragment(s)
	343	809
HindIII		A'AGCT_T (0 Err) - 2 Fragment(s)
	135	1017
NciI		CC's_GG (0 Err) - 2 Fragment(s)
	196	956
NcoI		C'CATG_G (0 Err) - 4 Fragment(s)
	60	135 288 669
PstI		C_TGCA'G (0 Err) - 3 Fragment(s)
	337	366 449
PvuII		CAG'CTG (0 Err) - 2 Fragment(s)
	570	582
Sall		G'TCGA_C (0 Err) - 2 Fragment(s)
	242	910

Linear Map of Sequence:

Figure 2-2

AccI
 241 gtcgacatggttctcatcccttgcggactcaggattcaaccatggtcagtgc
 cagctgtaccaagtagagggaaacgaaacctgagtcgtaacgttggaccacgtcaeg
 NcoI
 301 tttggccatatcagcggtggccacatcaaccctgcaggactgtggccatgggtgcacc
 aaaccggtatagtcgcaccggtagttggacgtcaactgacaccgttaccacacgtgg
 PstI NcoI
 361 aggaagatcagcatcgccaagtctgtcttctacatcgccaggccact
 tccttctagtcgtacgggtcagacagaagatgttagcgtcgggtcacggacccggtag
 421 attggagcaggaatcctctatctggcacacctcccagggtggggaggcctggagtc
 taacctcgcccttaggagatagaccaggactgtgggggtcacaccaccctccggaccctcag
 NcoI
 481 accatggttcatggaaatcttaccgctggtcatggtctcctgggtgatggataatcaca
 tggtaccaagtacctttagaattggcgaccaggactaccaggaccaactcaactattgt
 Pvull
 541 ttcaattggttactatcttgccaggctgtgatccaaacggactgttcactggc
 aaagttaaccacaaatgtatggcgacactaaagggttgcctgactacagtggacc
 601 tcaatagcttagcaattggattttctgttgcattggacattttgcattcaattat
 agttatcgaaatcgtaacctaagacaacgttaacctgttaataaaacgttagttaata
 PstI
 661 actgggccaggcatgaatccccccgatccttggacactgcaggatcatggaaattgg
 tgaccacggctgtacttagggcggttagggaaacctggacgtcaatagtacccttaacc
 Apal
 721 gaaaaccattggatataattgggtggccatcataggagctgtcctcgctggccctt
 ctggtaacctatataacccaaacccggtagtacccctcgacaggacggaccacccggaa
 EcoRI
 781 tatgagtagtcttctgtccagatgttgcattcaacgtcgatggaaacgttcggaaatcg
 atactcatacagaagacaggctacaacttaagttgcagaaaaattctcgaaatcg
 840

Figure 2-3

841	aaagctgcccaaaaaggaaagctacatggaggtggaggacaacaggagtccaggta tttcgacgggtcgtttccatgtacacctccaccctgttgcctcagtcatt	900
	NciI	
901	gagacggatgacctgattctaaaacacctggagtggtgcattgacgttgcacgggg ctctgcctactggactaagatTTGGACCTCACCACGTACACTAACTGCAACTGGCCCT	960
961	gaggagaagaaggggaaaagccaatctggagaggattgtcttcagtatgactagaagat ctccctttttcccccttctggtagaccttcataacagaagtcatactgtatcttcta	1020
1021	cgcactgaaagcagacaagactccttagaactgtcctcagatttcctccaccattaaag gcgtgactttcgctgttctgaggaaatcttgcacaggagtctaaaggaaagggtggtaattc	1080
1081	gaaacagatttgtataaattagaaatgtgcagggttggatgtcatattactca cttgcataacaatattatcacacgtccaaacaacaaagtacagtataatgagt	1140
1141	gtctaaacaata (SEQ ID NO:1) cagatttgttat	1200

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Citation

Algorithm Citation:

Not given.

Program Citation:

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